

GenCore version 5.1.6
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OM protein - protein search, using sw model

Run On: June 25, 2003, 14:42:11 ; Search time 11.5302 Seconds
(without alignments)
283.251 Million cell updates/sec

Title: US-09-622-613b-21

Perfect score: 605
Sequence: 1 MONNAFTFOKHIMTPIICN.....ICVCKCNGPYHFAIGRCRP 111

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database :

1: Issued_Patents_AA.*
2: /cgn2_6/ptodata/1/1aa/5A.COMB.pep.*
3: /cgn2_6/ptodata/1/1aa/5B.COMB.pep.*
4: /cgn2_6/ptodata/1/1aa/6A.COMB.pep.*
5: /cgn2_6/ptodata/1/1aa/6B.COMB.pep.*
6: /cgn2_6/ptodata/1/1aa/Backfile1.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	582.5	96.3	111	2	US-08-891-848-12
2	582.5	96.3	111	3	US-08-875-811-8
3	280.5	46.4	104	1	US-08-467-955-2
4	275.5	45.5	105	3	US-08-875-811-39
5	275.5	45.5	355	3	US-08-875-811-41
6	275.5	45.5	358	3	US-08-875-811-51
7	273.5	45.2	104	4	US-09-394-268-1
8	273.5	45.2	104	4	US-09-394-268-2
9	273.5	45.2	104	4	US-09-687-748-1
10	273.5	45.2	104	4	US-09-687-748-2
11	273.5	45.2	112	3	US-08-875-811-32
12	273.5	45.2	129	3	US-08-875-811-63
13	273.5	45.2	251	3	US-08-875-811-59
14	273.5	45.2	254	3	US-08-875-811-61
15	273.5	45.2	355	3	US-08-875-811-49
16	273.5	45.2	355	3	US-08-875-811-57
17	273.5	45.2	355	3	US-08-875-811-64
18	273.5	45.2	366	3	US-08-875-811-55
19	273.5	45.2	379	3	US-08-875-811-43
20	270.5	44.7	104	1	US-08-283-971-1
21	270.5	44.7	104	1	US-07-921-619-1
22	270.5	44.7	104	1	US-08-467-955-1
23	270.5	44.7	104	2	US-08-891-848-13
24	268.5	44.4	104	3	US-08-875-811-1
25	268.5	44.4	104	4	US-09-071-672-1
26	268.5	44.4	105	3	US-08-875-811-26
27	268.5	44.4	106	3	US-08-875-811-28

28	268.5	44.4	107	3	US-08-875-811-30	Sequence 30, Appl
29	267.5	44.2	105	3	US-08-875-811-24	Sequence 24, Appl
30	264.5	43.7	358	3	US-08-875-811-45	Sequence 45, Appl
31	264.5	43.7	365	3	US-08-875-811-53	Sequence 53, Appl
32	246.5	40.7	107	3	US-08-875-811-20	Sequence 20, Appl
33	235.5	38.9	111	3	US-08-875-811-22	Sequence 22, Appl
34	232	38.3	114	4	US-09-223-118-3	Sequence 47, Appl
35	228.5	37.8	360	3	US-08-875-811-47	Sequence 2, Appl
36	223	36.9	114	4	US-09-223-118-2	Sequence 1, Appl
37	222	36.7	114	4	US-09-223-118-1	Sequence 4, Appl
38	221	36.5	114	4	US-09-223-118-4	Sequence 3, Appl
39	202	33.4	83	3	US-08-875-811-2	Sequence 2, Appl
40	202	33.4	83	4	US-09-071-672-3	Sequence 2, Appl
41	161	26.6	169	1	US-08-441-629-2	Sequence 2, Appl
42	161	26.6	169	3	US-08-776-207-2	Sequence 2, Appl
43	161	26.6	169	4	US-09-507-773-2	Sequence 2, Appl
44	161	26.6	169	5	PCR-US95-09172-2	Sequence 2, Appl
45	123.5	20.4	125	6	5171845-2	Patent No. 5171845

ALIGNMENTS

RESULT 1
US-08-891-848-12
Sequence 12, Application US/08891848
Patent No. 5955073
GENERAL INFORMATION:
APPLICANT: Rybak, Susanna M.
APPLICANT: Youle, Richard J.
APPLICANT: Newton, Dianne L.
APPLICANT: Nicholls, Peter J.
TITLE OF INVENTION: Selective RNase Cytotoxic Reagents
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/891,848
FILING DATE: No. 5955073 yet assigned
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/125,462
FILING DATE: 22-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/014,082
FILING DATE: 04-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/779,195
FILING DATE: 22-OCT-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/510,696
FILING DATE: 20-APR-1990
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 015280-110310US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 111 amino acids
TYPE: amino acid

MOLECULE TYPE: protein
HYPOTHETICAL: N
ANTI-SENSE: N
FRAGMENT TYPE: N-termina
ORIGINAL SOURCE:
ORGANISM: Rana pipiens

DEVELOPMENTAL STAGE: Oocyte
US-08-467-955-2

Query Match	46.48;	Score 280.5;	DB 1;	Length 104;
Best Local Similarity	48.68;	Pred. No. 2.8e-25;		
Matches 54; Conservative 17; Mismatches 31; Indels 9; Gaps 4;				

RESULT 4
US-08-875-811-39
; Sequence 39, Application US/08875811
Detect No. 6045702

Query Match	45.5%	Score 275.5	DB 3	Length 105
Best Local Similarity	48.2%	Pred. No. 1.1e-24		
Matches 54	Conservative 17	Mismatches 32	Indels 9	Gaps 4

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Db      57 NVLTSEFYLSDC--NVTSRPCCKXIKLKSTNKFVCVTGCENQAFVHFVNGSC 105
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RESULT 5
US-08-875-811-41
; Sequence 41, Application US/08875811
Date: 08/04/2008 10:45:03

Query Match	45.5%	Score 275.5	DB 3	Length 355
Best Local Similarity	48.2%	Pred. No. 4.7e-24		
Matches 54	Conservative 17	Mismatches 32	Indels 9	Gaps 4

	Best Local Similarity	48.2%	Pred. No. 1.le-24	
Matches	54;	Conservative 17;	Mismatches 32;	Indels 9; Gaps 4
QY	1	MONATFOCKHIINT-PIICNTITLDNNIYVGGOCRVNFTFISSATVAKCTGYI-NL	58	
Db	1	MEDTLTFRKHTTNBDDCDINIMSTNF---HCKDKTFTIYSREPKAKCTGKGIASK	56	
QY	59	NVLSTTRQLNCTRTSTITPRCPYSSRTETNYICVCKENQVPHVAGIGRC	110	

APPLICANT: Wlodawer, Alexander
TITLE OF INVENTION: Recombinant Ribonuclease Proteins
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875.811
FILING DATE: 19-FEB-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/011.800
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Paris, Susan K.
REGISTRATION NUMBER: 41.739
REFERENCE/DOCKET NUMBER: 015280-244100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 358 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-875-811-51

Query Match: 45.5%; Score 275.5; DB 3; Length 358;
Best Local Similarity 48.2%; Pred. No. 4.8e-24;
Matches 54; Conservative 17; Mismatches 32; Indels 9; Gaps 4;
QY 1 QNMATFOQKHINT-PIICNTILDNNIYIGGCKRVNFIISATTVKAICGVI-NL 58
DB 1 MEDLFTQKHINTTRVDCCDNIINSTLNF---HCKDKNTFTYSPPEVKAICGIASK 56
QY 59 NVLSTTRQNLCTRTSITPRCPYSSRTETNYICVKCENQYPVHFGAGRC 110
DB 57 NVLTSEFYLSDC---NVTSRPKYKLLKSTNKFCTVCENQAPVHFGVGS 105
RESULT 7
US-09-394-268-1
Sequence 1, Application US/09394268
Patent No. 6175003
GENERAL INFORMATION:
APPLICANT: Saxena, Shalendra K
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING RIBONUCLEASES AND METHODS OF
FILE REFERENCE: 5013
CURRENT APPLICATION NUMBER: US/09/394,268
CURRENT FILING DATE: 1999-09-10
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 104
TYPE: PRN
ORGANISM: Rana pipiens
US-09-394-268-1
Query Match: 45.2%; Score 273.5; DB 4; Length 104;

Best Local Similarity 48.6%; Pred. No. 1.8e-24;
Matches 54; Conservative 16; Mismatches 32; Indels 9; Gaps 4;
QY 2 QNMATFOQKHINT-PIICNTILDNNIYIGGCKRVNFIISATTVKAICGVI-NLN 59
DB 1 QDMLTFOQKHINTTRVDCCDNIINSTLNF---HCKDKNTFTYSPPEVKAICGIASK 56
QY 60 NVLSTTRQNLCTRTSITPRCPYSSRTETNYICVKCENQYPVHFGAGRC 110
DB 57 NVLTSEFYLSDC---NVTSRPKYKLLKSTNKFCTVCENQAPVHFGVGS 104

RESULT 8
US-09-394-268-2
Sequence 2, Application US/09394268
Patent No. 6175003
GENERAL INFORMATION:
APPLICANT: Saxena, Shalendra K
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING RIBONUCLEASES AND METHODS OF
FILE REFERENCE: 5013
CURRENT APPLICATION NUMBER: US/09/394,268
CURRENT FILING DATE: 1999-09-10
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
LENGTH: 104
TYPE: PRN
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: SEQ ID NO:1 with Leu at position 23 and Cys at
OTHER INFORMATION: position 72
US-09-394-268-2

Query Match: 45.2%; Score 273.5; DB 4; Length 104;
Best Local Similarity 49.5%; Pred. No. 1.8e-24;
Matches 55; Conservative 15; Mismatches 32; Indels 9; Gaps 4;

QY 2 QNMATFOQKHINT-PIICNTILDNNIYIGGCKRVNFIISATTVKAICGVI-NLN 59
DB 1 QDMLTFOQKHINTTRVDCCDNIINSTLNF---HCKDKNTFTYSPPEVKAICGIASK 56
QY 60 NVLSTTRQNLCTRTSITPRCPYSSRTETNYICVKCENQYPVHFGAGRC 110
DB 57 NVLTSEFYLSDC---NVTSRPKYKLLKSTNKFCTVCENQAPVHFGVGS 104

RESULT 9
US-09-687-748-1
Sequence 1, Application US/09687748
Patent No. 6423515
GENERAL INFORMATION:
APPLICANT: Saxena, Shalendra K
TITLE OF INVENTION: METHODS OF MAKING NUCLEIC ACIDS ENCODING RIBONUCLEASES
FILE REFERENCE: 5013 US 01
CURRENT APPLICATION NUMBER: US/09/687,748
CURRENT FILING DATE: 2000-10-14
PRIOR APPLICATION NUMBER: 09/394,268
PRIOR FILING DATE: 1999-09-10
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1
LENGTH: 104
TYPE: PRN
ORGANISM: Rana pipiens
US-09-687-748-1

Query Match: 45.2%; Score 273.5; DB 4; Length 104;
Best Local Similarity 48.6%; Pred. No. 1.8e-24;
Matches 54; Conservative 16; Mismatches 32; Indels 9; Gaps 4;
QY 2 QNMATFOQKHINT-PIICNTILDNNIYIGGCKRVNFIISATTVKAICGVI-NLN 59
DB 1 QDMLTFOQKHINTTRVDCCDNIINSTLNF---HCKDKNTFTYSPPEVKAICGIASK 56

REGISTRATION NUMBER: 41/139
REFERENCE/DOCKET NUMBER: 015280-244100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 63:
SEQUENCE CHARACTERISTICS:

LENGTH: 129 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-08-875-811-63

Query Match 45.2%; Score 273.5; DB 3; Length 129;
 Best Local Similarity 48.6%; Pred. No. 2.3e-24;

Matches 54; Conservative 16; Mismatches 32; Indels 9; Gaps 4;

QY 2 ONMATEFOOKHIINT-PIICNTIILDNNIYVGGCKRVNFIISATTYKATGTVI-NLN 59
 Db 26 QDLTFOKKHITRTVDVCDNIMSTNLF---HCKDKNTFIYSRPEPVAKICKGIASKN 81
 QY 60 VLSTTFQOLNTCTRTSITPRCPYSSRTETNYICVGCENQYVHFAIGIRC 110
 Db 82 VLSTTFEYLSDC---NVTSRPKYKIKKSTNKFVCYCENQAPVHFGVASC 129

RESULT 13

US-08-875-811-59
 Sequence 59, Application US/08875811
 Patent No. 6045793

GENERAL INFORMATION:

APPLICANT: Rybak, Susanna M.
 APPLICANT: Newton, Dianne L.
 APPLICANT: Bogue, Luis
 APPLICANT: Wlodawer, Alexander
 TITLE OF INVENTION: Recombinant Ribonuclease Proteins
 NUMBER OF SEQUENCES: 64
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Townsend and Townsend and Crew LLP
 STREET: Two Embarcadero Center, Eighth Floor
 CITY: San Francisco
 STATE: California
 COUNTRY: USA
 ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patentin Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/875,811
 FILING DATE: 19-FEB-1998
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: WO PCT/US97/02588
 FILING DATE: 19-FEB-1997

PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 60/011,800
 FILING DATE: 21-FEB-1996
 ATTORNEY/AGENT INFORMATION:
 NAME: Faris, Susan K.
 REGISTRATION NUMBER: 41,739

REFERENCE/DOCKET NUMBER: 015280-244100US
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (415) 576-0200
 TELEFAX: (415) 576-0300

INFORMATION FOR SEQ ID NO: 59:

SEQUENCE CHARACTERISTICS:
 LENGTH: 251 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-08-875-811-59

Query Match 45.2%; Score 273.5; DB 3; Length 251;
 Best Local Similarity 48.2%; Pred. No. 5.3e-24;

Matches 54; Conservative 16; Mismatches 33; Indels 9; Gaps 4;

QY 1 MONMATEFOOKHIINT-PIICNTIILDNNIYVGGCKRVNFIISATTYKATGTVI-NL 58
 Db 1 MSDWLTFOKKHITRTVDVCDNIMSTNLF---HCKDKNTFIYSRPEPVAKICKGIASKN 56

Db 147 MSDWLTFOKKHITRTVDVCDNIMSTNLF---HCKDKNTFIYSRPEPVAKICKGIASKN 202
 QY 59 NVLSTTFQOLNTCTRTSITPRCPYSSRTETNYICVGCENQYVHFAIGIRC 110
 Db 203 NVLSTTFEYLSDC---NVTSRPKYKIKKSTNKFVCYCENQAPVHFGVASC 251

RESULT 14

US-08-875-811-61
 Sequence 61, Application US/08875811
 Patent No. 6045793

GENERAL INFORMATION:

APPLICANT: Rybak, Susanna M.
 APPLICANT: Newton, Dianne L.
 APPLICANT: Bogue, Luis
 APPLICANT: Wlodawer, Alexander
 TITLE OF INVENTION: Recombinant Ribonuclease Proteins
 NUMBER OF SEQUENCES: 64
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Townsend and Townsend and Crew LLP
 STREET: Two Embarcadero Center, Eighth Floor
 CITY: San Francisco
 STATE: California
 COUNTRY: USA
 ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patentin Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/875,811
 FILING DATE: 19-FEB-1998
 CLASSIFICATION: 435
 PRIOR APPLICATION DATA:
 APPLICATION NUMBER: WO PCT/US97/02588
 FILING DATE: 19-FEB-1997

PRIOR APPLICATION DATA:
 APPLICATION NUMBER: US 60/011,800
 FILING DATE: 21-FEB-1996
 ATTORNEY/AGENT INFORMATION:
 NAME: Faris, Susan K.
 REGISTRATION NUMBER: 41,739

REFERENCE/DOCKET NUMBER: 015280-244100US
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (415) 576-0200
 TELEFAX: (415) 576-0300

INFORMATION FOR SEQ ID NO: 61:

SEQUENCE CHARACTERISTICS:
 LENGTH: 254 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-08-875-811-61

Query Match 45.2%; Score 273.5; DB 3; Length 254;
 Best Local Similarity 48.2%; Pred. No. 5.3e-24;

Matches 54; Conservative 16; Mismatches 33; Indels 9; Gaps 4;

QY 1 MONMATEFOOKHIINT-PIICNTIILDNNIYVGGCKRVNFIISATTYKATGTVI-NL 58
 Db 1 MSDWLTFOKKHITRTVDVCDNIMSTNLF---HCKDKNTFIYSRPEPVAKICKGIASKN 56
 QY 59 NVLSTTFQOLNTCTRTSITPRCPYSSRTETNYICVGCENQYVHFAIGIRC 110
 Db 57 NVLSTTFEYLSDC---NVTSRPKYKIKKSTNKFVCYCENQAPVHFGVASC 105

RESULT 15

US-08-875-811-49
 Sequence 49, Application US/08875811
 Patent No. 6045793

GENERAL INFORMATION:

APPLICANT: Rybak, Susanna M.
APPLICANT: Newton, Dianne L.
APPLICANT: Bogue, Julius
APPLICANT: Widawer, Alexander
TITLE OF INVENTION: Recombinant Ribonuclease, Proteins
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3634
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/875,811
FILING DATE: 19-FEB-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US97/02588
FILING DATE: 19-FEB-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/011,800
FILING DATE: 21-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: Paris, Susan K.
REGISTRATION NUMBER: 41,739
REFERENCE/DCKET NUMBER: 015280-244100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
LENGTH: 355 amino acids
TYPE: amine acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-875-811-49

Query Match	Similarity	45.28	Score 273.5	DB 3	Length 355
Best Local	Similarity	48.28	Pred. No. 8e-24		
Matches	Conservative	16	Mismatches	33	Indels 9; Gaps 4
Oy	1	MQNNATFOCKHILNM-PIICMTIIDNNIIYIGGCKRNTFIISATYKATCGVI-NL	58		
Db	251	MSDMITFOCKHITINRDVCDNIMSTNF---HCKDNFTFYSSRPEPKACIKGIASK	306		
Oy	59	NVLSTTRQLNCTRTSTTPRCPYSSRPTENIVCKENQVPMFAGIGRC	110		
Db	307	NVLSTSEFYSDC---NVTSRCKKLLKSTNKECVCTENQAPVHFVGVGSC	355		

Search completed: June 25, 2003, 15:00:31
Job time : 20.5302 secs

